

0304/0150  
1130

OIPE

#3

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/897,107DATE: 11/20/2001  
TIME: 10:09:44Input Set : A:\ES.txt  
Output Set: N:\CRF3\11202001\I897107.raw**ENTERED**

3 <110> APPLICANT: YAMAGISHI, Akihiko  
 5 <120> TITLE OF INVENTION: METHOD FOR IMPROVING THERMOSTABILITY OF PROTEINS, PROTEINS  
 HAVING  
 6 THERMOSTABILITY IMPROVED BY THE METHOD AND NUCLEIC ACIDS ENCODING THE PROTEINS  
 8 <130> FILE REFERENCE: 210383US0  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/897,107  
 11 <141> CURRENT FILING DATE: 2001-07-03  
 13 <150> PRIOR APPLICATION NUMBER: JP2000-201920  
 14 <151> PRIOR FILING DATE: 2000-07-04  
 16 <150> PRIOR APPLICATION NUMBER: JP2001-164332  
 17 <151> PRIOR FILING DATE: 2001-05-31  
 19 <160> NUMBER OF SEQ ID NOS: 104  
 21 <170> SOFTWARE: PatentIn version 3.1  
 23 <210> SEQ ID NO: 1  
 24 <211> LENGTH: 9  
 25 <212> TYPE: PRT  
 26 <213> ORGANISM: Sulfolobus sp.  
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 31 1 5  
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 36 <212> TYPE: PRT  
 37 <213> ORGANISM: Sulfolobus sp.  
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 47 <212> TYPE: PRT  
 48 <213> ORGANISM: Sulfolobus sp.  
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 53 1 5  
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 57 <211> LENGTH: 6  
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 59 <213> ORGANISM: Sulfolobus sp.  
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 64 1 5  
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 68 <211> LENGTH: 9  
 69 <212> TYPE: PRT  
 70 <213> ORGANISM: Thermus thermophilus  
 72 <400> SEQUENCE: 5  
 74 Gln Asp Leu Phe Ala Asn Leu Arg Pro  
 75 1 5

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Input Set : A:\ES.txt

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86 1 5  
89 <210> SEQ ID NO: 7  
90 <211> LENGTH: 8  
91 <212> TYPE: PRT  
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94 <400> SEQUENCE: 7  
96 Val His Gly Ser Ala Pro Asp Ile  
97 1 5  
100 <210> SEQ ID NO: 8  
101 <211> LENGTH: 6  
102 <212> TYPE: PRT  
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108 1 5  
111 <210> SEQ ID NO: 9  
112 <211> LENGTH: 9  
113 <212> TYPE: PRT  
114 <213> ORGANISM: Bacillus subtilis  
116 <400> SEQUENCE: 9  
118 Leu Asp Leu Phe Ala Asn Leu Arg Pro  
119 1 5  
122 <210> SEQ ID NO: 10  
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125 <213> ORGANISM: Bacillus subtilis  
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129 Val Ile Arg Glu Gly Phe Lys Met Ala  
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149 <400> SEQUENCE: 12  
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152 1 5

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Input Set : A:\ES.txt

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163 1 5  
166 <210> SEQ ID NO: 14  
167 <211> LENGTH: 9  
168 <212> TYPE: PRT  
169 <213> ORGANISM: Escherichia coli  
171 <400> SEQUENCE: 14  
173 Ile Ala Arg Ile Ala Phe Glu Ser Ala  
174 1 5  
177 <210> SEQ ID NO: 15  
178 <211> LENGTH: 8  
179 <212> TYPE: PRT  
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184 Ala Gly Gly Ser Ala Pro Asp Ile  
185 1 5  
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189 <211> LENGTH: 6  
190 <212> TYPE: PRT  
191 <213> ORGANISM: Escherichia coli  
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195 Leu Leu Leu Arg Tyr Ser  
196 1 5  
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200 <211> LENGTH: 9  
201 <212> TYPE: PRT  
202 <213> ORGANISM: Agrobacterium tumefaciens  
204 <400> SEQUENCE: 17  
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207 1 5  
210 <210> SEQ ID NO: 18  
211 <211> LENGTH: 9  
212 <212> TYPE: PRT  
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221 <210> SEQ ID NO: 19  
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223 <212> TYPE: PRT  
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226 <400> SEQUENCE: 19  
228 Val His Gly Ser Ala Pro Asp Ile  
229 1 5

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Input Set : A:\ES.txt  
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240 1 5
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246 <213> ORGANISM: Saccharomyces cerevisiae
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250 Leu Gln Leu Tyr Ala Asn Leu Arg Pro
251 1 5
254 <210> SEQ ID NO: 22
255 <211> LENGTH: 9
256 <212> TYPE: PRT
257 <213> ORGANISM: Saccharomyces cerevisiae
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262 1 5
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267 <212> TYPE: PRT
268 <213> ORGANISM: Saccharomyces cerevisiae
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272 Cys His Gly Ser Ala Pro Asp Leu
273 1 5
276 <210> SEQ ID NO: 24
277 <211> LENGTH: 6
278 <212> TYPE: PRT
279 <213> ORGANISM: Saccharomyces cerevisiae
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284 1 5
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289 <212> TYPE: PRT
290 <213> ORGANISM: Neurospora crassa
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294 Leu Gly Thr Tyr Gly Asn Leu Arg Pro
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299 <211> LENGTH: 9
300 <212> TYPE: PRT
301 <213> ORGANISM: Neurospora crassa
303 <400> SEQUENCE: 26
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306 1 5
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311 <212> TYPE: PRT
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317 1 5
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321 <211> LENGTH: 6
322 <212> TYPE: PRT
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325 <400> SEQUENCE: 28
327 Met Met Leu Arg Tyr Ser
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332 <211> LENGTH: 9
333 <212> TYPE: PRT
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342 <210> SEQ ID NO: 30
343 <211> LENGTH: 9
344 <212> TYPE: PRT
345 <213> ORGANISM: Bos taurus
347 <400> SEQUENCE: 30
349 Val Ile Arg Tyr Ala Phe Glu Tyr Ala
350 1 5
353 <210> SEQ ID NO: 31
354 <211> LENGTH: 8
355 <212> TYPE: PRT
356 <213> ORGANISM: Saccharomyces cerevisiae
358 <400> SEQUENCE: 31
360 Val His Gly Ser Ala Pro Asp Ile
361 1 5
364 <210> SEQ ID NO: 32
365 <211> LENGTH: 6
366 <212> TYPE: PRT
367 <213> ORGANISM: Saccharomyces cerevisiae
369 <400> SEQUENCE: 32
371 Met Met Leu Asn His Met
372 1 5
375 <210> SEQ ID NO: 33
376 <211> LENGTH: 9
377 <212> TYPE: PRT
378 <213> ORGANISM: Bos taurus
380 <400> SEQUENCE: 33
382 Phe Asp Leu Tyr Ala Asn Val Arg Pro
383 1 5

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→ Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

## VERIFICATION SUMMARY

DATE: 11/20/2001

PATENT APPLICATION: US/09/897,107

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Input Set : A:\ES.txt

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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:529 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45  
L:561 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46  
L:607 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48